

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/846,687B
Source: FW/b
Date Processed by STIC: 10/16/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

09/846,687B

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 ☐ Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 ☐ Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

- 4 ☒ Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

- 5 ☐ Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 ☐ PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

- 7 ☐ Skipped Sequences
 (OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

- 8 ☐ Skipped Sequences
 (NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 ☐ Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 ☒ Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

- 11 ☐ Use of <220> Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

- 12 ☐ PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 ☐ Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFW16

RAW SEQUENCE LISTING

DATE: 10/16/2006

PATENT APPLICATION: US/09/846,687B

TIME: 12:16:26

Input Set : A:\PTO.SS.txt

Output Set: N:\CRF4\10162006\I846687B.raw

3 <110> APPLICANT: Lorrence H. Green
 5 <120> TITLE OF INVENTION: A Method and Vaccine for the Prevention of AIDS.
 W--> 0 <130> FILE REFERENCE:
 7 <140> CURRENT APPLICATION NUMBER: US 09/846,687B
 9 <141> CURRENT FILING DATE: 2001-05-01

*see item 4 on Euro
summary
sheet*

11 <160> NUMBER OF SEQ ID NOS: 3
 13 <210> SEQ ID NO: 1
 14 <211> LENGTH: 45
 15 <212> TYPE: PRT
 16 <213> ORGANISM: Artificial Sequence
 W--> 17 <220> FEATURE: Chemically Synthesized
 W--> 19 <223> OTHER INFORMATION: (C)
 W--> 19 <400> 1

*move this to <223> line.
<220> is a header only.*

*Does Not Comply
Corrected Diskette Needed
never
has a
response.*

21 Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile Leu
 22 1 5 10 15
 24 Gly Leu Val Leu Pro Leu Leu Val Met Val Ile Cys Tyr Ser Gly Ile
 25 20 25 30
 27 Leu Lys Thr Leu Leu Arg Cys Arg Asn Glu Lys Lys Arg
 28 35 40 45

30 <210> SEQ ID NO: 2
 31 <211> LENGTH: 45
 32 <212> TYPE: PRT
 33 <213> ORGANISM: Artificial Sequence
 W--> 34 <220> FEATURE: Chemically Synthesized
 W--> 36 <223> OTHER INFORMATION: (C)
 W--> 36 <400> 2

move down

*Insert <220> wherever
<221>, <222>, or <223>
is present.
e.g. <220>
<223> chemically
Synthesized*

38 Tyr Thr Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile Leu
 39 1 5 10 15
 41 Gly Leu Val Leu Pro Leu Leu Val Met Val Ile Cys Tyr Ser Gly Ile
 42 20 25 30
 44 Leu Lys Thr Leu Leu Arg Cys Arg Asn Glu Lys Lys Arg
 45 35 40 45

47 <210> SEQ ID NO: 3
 48 <211> LENGTH: 45
 49 <212> TYPE: PRT
 50 <213> ORGANISM: Polypeptide sp.
 W--> 51 <220> FEATURE: Chemically Synthesized
 53 <400> SEQUENCE: 3

*invalid response. See item 10 on Euro summary
sheet.*

<223> move to <223> line

55 Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile Leu
 56 1 5 10 15
 58 Gly Leu Val Leu Pro Leu Leu Val Cys Val Ile Cys Tyr Ser Gly Ile
 59 20 25 30
 61 Leu Lys Thr Leu Leu Arg Cys Arg Asn Glu Lys Lys Arg

RAW SEQUENCE LISTING

DATE: 10/16/2006

PATENT APPLICATION: US/09/846,687B

TIME: 12:16:26

Input Set : A:\PTO.SS.txt

Output Set: N:\CRF4\10162006\I846687B.raw

62

35

40

45

3-

VERIFICATION SUMMARY

DATE: 10/16/2006

PATENT APPLICATION: US/09/846,687B

TIME: 12:16:27

Input Set : A:\PTO.SS.txt

Output Set: N:\CRF4\10162006\I846687B.raw

L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE
L:17 M:283 W: Missing Blank Line separator, <220> field identifier
L:17 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:19 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:1, <213>
ORGANISM:Artificial Sequence
L:19 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:19
L:34 M:283 W: Missing Blank Line separator, <220> field identifier
L:34 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:36 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:2, <213>
ORGANISM:Artificial Sequence
L:36 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:2,Line#:36
L:51 M:283 W: Missing Blank Line separator, <220> field identifier
L:51 M:256 W: Invalid Numeric Header Field, <220> has non-blank data